Overlap of functional cREs with (epi)genomic features All_DNase All_DNase All_cCREs All_cCREs K562_ATAC K562_ATAC K562 DNase K562_DNase K562_cCREs K562_H3K4me1 K562_cCREs-K562_H3K4me3 K562_H3K4me1 Feature K562_H3K27ac K562 H3K4me3-K562 CTCF K562_H3K27ac-K562_EP300 K562_CTCF K562_POLR2A K562_RAMPAGE K562_EP300 K562_CAGE K562_POLR2A-K562 GROCAP K562 H3K9me3 K562_H3K9me3 K562_H3K27me3 K562_H3K27me3 25 50 75 100 -5.0 -2.5 0.02.5 Percent of Significant Eleme log2(Odds Ratio)